766.21 CIP

PATENT PPLICATION BOX/SEq.

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

an re Application of:)
TETSUYOSHI ISHIWATA et a	Examiner: N/Y/A
Application No: 09/730,559	Group Art Unit: N/Y/A
Filed: December 7, 2000	;)
For: IgA NEPHROPATHY-GENES	EELATED) August 3, 2001

Commissioner for Patents Washington, D.C. 20231

RESPONSE TO NOTICE TO COMPLY WITH REQUIREMENTS and SUBMISSION OF SUBSTITUTE COMPUTER READABLE FORM AND PAPER COPY

Sir:

This is in response to the Office Action mailed July 10, 2001 (copy attached). Applicants submit herewith a substitute computer readable form under 37 C.F.R. § 1.821(e). The content of the computer readable form and the Paper Copy of the Sequence Listing filed herewith are the same. No new matter has been added.

Applicants' undersigned attorney may be reached in our New York office by telephone at (212) 218-2100. All correspondence should continue to be directed to our below listed address.

Respectfully submitted,

Attorney for Applicants

Lawrence S. Perry Registration No. 31,865

FITZPATRICK, CELLA, HARPER & SCINTO 30 Rockefeller Plaza
New York, New York 10112-3801
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FILING/RECEIPT DATE FIRST NAMED APPLICANT ATTORNEY I

Office

09/730,559

12/07/2000

Tetsuyoshi Ishiwata

ATTORNEY DOCKET NUMBER
766.21 CIP

05514 FITZPATRICK CELLA HARPER & SCINTO 30 ROCKEFELLER PLAZA NEW YORK, NY 10112 CONFIRMATION NO. 9523
FORMALITIES LETTER

OC000000006275848

Date Mailed: 07/10/2001

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant is given **TWO MONTHS FROM THE DATE OF THIS NOTICE** within which to file the items indicated below to avoid abandonment. Extensions of time may be obtained under the provisions of 37 CFR 1.136(a).

• A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing." Applicant must provide a substitute computer readable form (CRF) copy of the "Sequence Listing" and a statement that the content of the sequence listing information recorded in computer readable form is identical to the written (on paper or compact disc) sequence listing and, where applicable, includes no new matter, as required by 37 CFR 1.821(e), 1.821(f), 1.821(g), 1.825(b), or 1.825(d).

For questions regarding compliance to these requirements, please contact:

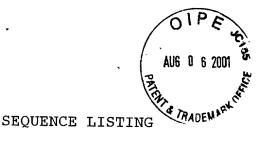
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PART 1 - ATTORNEY/APPLICANT COPY



<110> ISHIWATA, TETSUYOSHI
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 KAWABATA, AYAKO
 NAKAGAWA, SATOSHI
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Ala Phe Cys Pro Glu Leu Thr Pro Phe Pro Asp Gln Gly Thr Ser Thr 55 60 65 70	
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gct gtg t Ala Val T												162
ctt gcc a Leu Ala T									_		_	210
gtg cca g Val Pro G		_	_		-					Ile		258
aca tac o		-	_	ı Thr				_	-			306
gca cct c Ala Pro 6			_									354
tca gca t Ser Ala E 85	_	_	_			_			_	_		402

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450

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105 110 115

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Thr Lys Tyr Asp Val Leu Pro Tyr Ser Ile Arg Val Leu Leu Glu Ala
35 40 45

gct gta cga aat tgt gat ggc ttt tta atg aag aag gaa gat gtt atg 192 Ala Val Arg Asn Cys Asp Gly Phe Leu Met Lys Lys Glu Asp Val Met 50 55 60

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	gat Asp						-	_							-	336
	gag Glu					_	_	_		-			_	-		384
	tta Leu 130										_		-			432
Pro 145	gga Gly	Gly	Gly	Asp	Leu 150	Gln	Lys	Ala	Gly	Lys 155	Leu	Ser	Pro	Leu	Lys 160	480
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	tgt Cys				_			_							_	576
_	att Ile						_	-				_				624
	gaa Glu 210		-							_	-	-			_	672
	cga Arg				_			_		_		_	_		-	720
	gtg Val	-								_	_					768

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														aat Asn		864
														gca Ala		912
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gtaaaatcac	aaaaggtaag	ttgttggaag	acaacaaaaa	agaattacta	tatctgatcc	180
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ctctctggca gtgctccctg caaatgtgtc ctttcaagaa aacaaaacct gcaagtggct
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tgtaatgtac catgacctta tcatgtgaag gacaaatggc tcttgtgctt attagatagc
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aaaacaagtt taaaactcaa aagaggatta ttctcaagtt atactacagt gaaaaaacat
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<213> Homo sapiens

<400> 34

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Ser Gly Leu Gly Ser Pro His Cys Phe Ser His Gln Asn Gly Glu Arg
20 25 30

Val Glu Arg Tyr Ser Arg Lys Val Phe Val Gly Gly Leu Pro Pro Asp 35 40 45

Ile Asp Glu Asp Glu Ile Thr Ala Ser Phe Arg Arg Phe Gly Pro Leu 50 55 60

Ile Val Asp Trp Pro His Lys Ala Glu Ser Lys Ser Tyr Phe Pro Pro 65 70 75 80

Lys Gly Tyr Ala Phe Leu Leu Phe Gln Asp Glu Ser Ser Val Gln Ala 85 90 95

Leu Ile Asp Ala Cys Ile Glu Glu Asp Gly Lys Leu Tyr Leu Cys Val 100 105 110

Ser Ser Pro Thr Ile Lys Asp Lys Pro Val Gln Ile Arg Pro Trp Asn 115 120 125

Leu Ser Asp Ser Asp Phe Val Met Asp Gly Ser Gln Pro Leu Asp Pro 130 135 140

Arg Lys Thr Ile Phe Val Gly Gly Val Pro Arg Pro Leu Arg Ala Val 145 150 155 160

Glu Leu Ala Met Val Met Asp Arg Leu Tyr Gly Gly Val Cys Tyr Ala 165 170 175

Gly Ile Asp Thr Asp Pro Glu Leu Lys Tyr Pro Lys Gly Ala Gly Arg 180 185 190

Val Ala Phe Ser Asn Gln Gln Ser Tyr Ile Ala Ala Ile Ser Ala Arq

Phe Val Gln Leu Gln His Gly Glu Ile Asp Lys Arg Val Ser Leu Ile 210 215 220

Leu His Phe Gly Lys Phe 225 230

<210> 35

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Met Gly Ser Asp Lys Arg Val Ser Arg Thr Glu Arg Ser Gly Arg Tyr 1 5 10 15

Gly Ser Ile Ile Asp Arg Asp Asp Arg Asp Glu Arg Glu Ser Arg Ser 20 25 30

Arg Arg Arg Asp Ser Asp Tyr Lys Arg Ser Ser Asp Asp Arg Gly 35 40 45

Asp Arg Tyr Asp Asp Tyr Arg Asp Tyr Asp Ser Pro Glu Arg Glu Arg 50 55 60

Glu Arg Arg Asn Ser Asp Arg Ser Glu Asp Gly Tyr His Ser Asp Gly 65 70 75 80

Asp Tyr Gly Glu His Asp Tyr Arg His Asp Ile Ser Asp Glu Arg Glu 85 90 95

Ser Lys Thr Ile Met Leu Arg Gly Leu Pro Ile Thr Ile Thr Glu Ser 100 105 110

Asp Ile Arg Glu Met Met Glu Ser Phe Glu Gly Pro Gln Pro Ala Asp 115 120 125

Val Arg Leu Met Lys Arg Lys Thr Gly Glu Ser Leu Leu Ser Ser 130 135 140

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<211> 104

<212> PRT

<213> Homo sapiens

<400> 36

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Cys Val Thr Ala Leu Gly Glu Glu Thr Gly Ala Trp Phe Pro Val Tyr
20 25 30

Leu Ser His Ala Ser Ser Pro Phe Ala Asp Leu Val Phe Cys Pro Phe 35 40 45

Ala Glu Ile Asn His Ser Gln Glu Tyr Asp Asn Met Arg Gly Pro Val
50 55 60

Ser Pro Pro Asn Lys Gln Phe Asn Leu Gly Val Ile Phe Gly Ile Pro 65 70 75 80

Asn Asn Cys Arg Phe Pro Thr Asp Asn Lys Ile Thr Glu Lys Gln Leu 85 90 95

Leu Gly Asn Val Leu Asn Tyr Pro 100

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<213> Homo sapiens

<400> 37

Met Asn His Pro Trp His Val Cys Phe Leu Phe Lys Val Leu Arg Tyr 1 5 10 15

Tyr Pro Thr Ala Pro Ile Leu Lys Trp Thr His Thr Val Ser Cys Ser 20 25 30

Trp Cys Arg Ser Val Leu Arg Glu Val Val Gly Asn Val Ser Leu Ser 35 40 45

Glu Asn Phe Thr Ile Ser Ala Phe Cys Pro Glu Leu Thr Pro Phe Pro 50 55 60

Asp Gln Gly Thr Ser Thr Met Ile Ser Phe Leu Glu Lys Phe Asn Lys 65 70 75 80

Ser Lys Arg Glu Arg Leu Glu Leu Met Leu His Phe Tyr Ser Val Leu 85 90 95

Ser Leu Glu Pro Ala Val Ala Glu His Trp Ser Gly Glu Phe Glu Lys 100 105 110

Trp Lys Val Gly Phe Phe His Pro Leu Lys Arg Glu Asp Gly Phe Phe 115 120 125

Thr Arg Thr Asp Ile 130

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<211> 133

<212> PRT

<213> Homo sapiens

<400> 38

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Tyr Pro Thr Ala Pro Ile Leu Lys Trp Thr His Thr Val Ser Cys Ser 20 25 30

Trp Cys Arg Ser Val Leu Arg Glu Val Val Gly Asn Val Ser Leu Ser 35 40 45

Glu Asn Phe Thr Ile Ser Ala Phe Cys Pro Glu Leu Thr Pro Phe Pro 50 55 60

Asp Gln Gly Thr Ser Thr Met Ile Ser Phe Leu Glu Lys Phe Asn Lys 65 70 75 80

Ser Lys Arg Glu Arg Leu Glu Leu Met Leu His Phe Tyr Ser Val Leu 85 90 95

Ser Leu Glu Pro Ala Phe Ala Glu His Trp Ser Gly Glu Phe Glu Lys 100 105 110

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Glu Lys Leu Leu Leu Ala Thr Gly Leu Asp Gly Ser Tyr Leu Leu Arg
20 25 30

Asp Ser Glu Ser Val Pro Gly Val Tyr Cys Leu Cys Val Leu Tyr His
35 40 45

Gly Tyr Ile Tyr Thr Tyr Arg Val Ser Gln Thr Glu Thr Gly Ser Trp
50 55 60

Ser Ala Glu Thr Ala Pro Gly Val His Lys Arg Tyr Phe Arg Lys Ile
65 70 75 80

Lys Asn Leu Ile Ser Ala Phe Gln Lys Pro Asp Gln Gly Ile Val Ile 85 90 95

Pro Leu Gln Tyr Pro Val Glu Lys Lys Ser Ser Ala Arg Ser Thr Gln
100 105 110

Gly Thr Thr Gly Ile Arg Glu Asp Pro Asp Val Cys Leu Lys Ala Pro 115 120 125

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Leu Asn Asp Ser Ser His Lys Lys Phe Phe Asp Val Ser Lys Leu Gly
20 25 30

Thr Lys Tyr Asp Val Leu Pro Tyr Ser Ile Arg Val Leu Leu Glu Ala 35 40 45

Ala Val Arg Asn Cys Asp Gly Phe Leu Met Lys Lys Glu Asp Val Met 50 55 60

Asn Ile Leu Asp Trp Lys Thr Lys Gln Ser Asn Val Glu Val Pro Phe 65 70 75 80

Phe Pro Ala Arg Val Leu Gln Asp Phe Thr Gly Ile Pro Ala Met 85 90 95

Val Asp Phe Ala Ala Met Arg Glu Ala Val Lys Thr Leu Gly Gly Asp 100 105 110

Pro Glu Lys Val His Pro Ala Cys Pro Thr Asp Leu Thr Val Asp His 115 120 125

Ser Leu Gln Ile Asp Phe Ser Lys Cys Ala Ile Gln Asn Ala Pro Asn 130 135 140

Pro Gly Gly Asp Leu Gln Lys Ala Gly Lys Leu Ser Pro Leu Lys 145 150 155 160

Val Gln Pro Lys Lys Leu Pro Cys Arg Gly Gln Thr Thr Cys Arg Gly
165 170 175

Ser Cys Asp Ser Gly Glu Leu Gly Arg Asn Ser Gly Thr Phe Ser Ser 180 185 190

Gln Ile Glu Asn Thr Pro Ile Leu Cys Pro Phe His Leu Gln Pro Val 195 200 205

Pro Glu Pro Glu Thr Val Leu Lys Asn Gln Glu Val Glu Phe Gly Arg 210 215 220

Asn Arg Glu Arg Leu Gln Phe Phe Lys Trp Ser Ser Arg Val Leu Lys 225 230 235 240

Asn Val Ala Val Ile Pro Pro Gly Thr Gly Met Ala His Gln Ile Asn 245 250 255

Leu Glu Tyr Leu Ser Arg Val Val Phe Glu Glu Lys Asp Leu Leu Phe 260 265 270

Pro Asp Ser Val Val Gly Thr Asp Ser His Ile Thr Met Val Asn Gly 275 280 285

Leu Gly Ile Leu Gly Trp Gly Val Gly Gly Ile Glu Thr Glu Ala Val

Met Leu Gly I 305	Leu Pro Val Ser Le 310	u Thr Leu Pro 315	Glu Val Val	Gly Cys 320
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cataatttaa ac	caaaatcaa ctaagatga	at ccaagttcca	cacaactgca	cttcaatatt 180
caagtcggtg tg	gaagatgcc tgactacto	gc gtcacaagat	tctgagctgt	cgtaaaaagc 240
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300

295

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290

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<213> Homo sapiens

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tatgtacaca gtgagaggat acttgtagag aacctagaat cttctctgaa tgtgacgggc 1	180
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<210> 70
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<400> 70
gagagagatg tcagagtcat tagc
                                                                        24
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<400> 71
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<400> 77 gctgaaacct aagctgaagg aagg	24
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<400> 78 gtccctcacc tcagatcaca cc	22

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gctatctacc tggcaggaaa agag
                                                                        24
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gagtttctta ctatgatctg gattc
                                                                        25
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gcaaaatgta ctcagcttca atcac
                                                                        25
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gcataacaag tgacagggtt agtta
                                                                        25
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<223> Description of Artificial Sequence: Synthetic DNA
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ggtgctcctt ccttacactg gt
                                                                        22
<210> 88
<211> 23
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gactacacat aaacccaccc cag
                                                                        23
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<400> 93
gacagcaacc taataacagc tgtc
                                                                         24
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gtcctaggca cttgtcacta qq
                                                                        22
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gaggggactt ccaagagtct ct
                                                                        22
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gacttcctaa ggcacactca gc	22
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gassaagaas coagecoace agec	24
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gaatgaacca gagccaggac ag	22

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gccttgtatg tatgcctgtg cc
                                                                         22
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aagagtccac caggccatgg a
                                                                        21
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<400> 106
taccttgtgt acttctagct gag
                                                                        23
<210> 107
<211> 17
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•	<210> 108 <211> 17 <212> DNA <213> Artificial Sequence	
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	<400> 109 gtttttttt tttttc	17
	<210> 110 <211> 18 <212> DNA <213> Artificial Sequence	
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	<400> 110 cagagtgatg gatatcaa	18

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atgaaagtgc cagtgtgcca tg
                                                                        22
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<223> Description of Artificial Sequence:Synthetic DNA
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cccatcacca tcttccagga gc
                                                                        22
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ttcaccacct tcttgatgtc atcata
                                                                        26
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<223> Description of Artificial Sequence: Synthetic Peptide
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<400> 114
Cys Pro Leu Lys Arg Glu Asp Gly Phe Phe Thr Arg Thr Asp Ile
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                   5
                                       10
                                                            15
<210> 115
<211> 16
<212> PRT
<213> Artificial Sequence
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<221> MOD RES
<222> (16)
<223> AMIDATION, GluAmide
<400> 115
Cys Ser Phe Leu Glu Lys Phe Asn Lys Ser Lys Arg Glu Arg Leu Xaa
  1
                   5
                                       10
                                                           15
<210> 116
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<221> MOD RES
<222> (15)
<223> AMIDATION, GlyAmide
<400> 116
Cys Ala Glu His Trp Ser Gly Glu Phe Glu Lys Trp Lys Val Xaa
  1
                                      10
                                                           15
<210> 117
<211> 16
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic Peptide
<400> 117
Cys Glu Ile Asp Lys Arg Val Ser Leu Ile Leu His Phe Gly Lys Phe
                   5
                                      10
                                                           15
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<210> 118
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<220>
<223> Description of Artificial Sequence: Synthetic Peptide
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Cys Arg Leu Met Lys Arg Lys Thr Gly Glu Ser Leu Leu Ser Ser
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                                                           15
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<211> 14
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Cys Thr Ser Ile Asp Val Val Leu Gly Ile Thr Lys Val Ser
<210> 120
<211> 16
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<222> (16)
<223> AMIDATION, LysAmide
<400> 120
Cys Ser Ala Glu Thr Ala Pro Gly Val His Lys Arg Tyr Phe Arg Xaa
                                      10
                                                           15
<210> 121
<211> 16
<212> PRT
<213> Artificial Sequence
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